

FIG. 1

ì

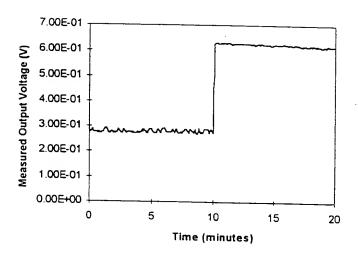


FIG. 2

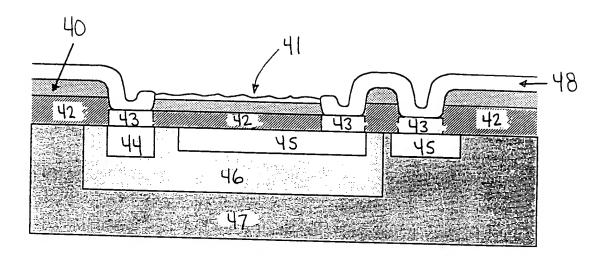


FIG. 3A

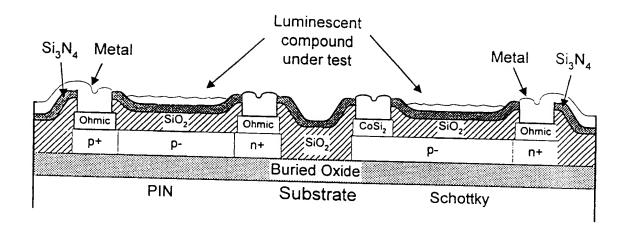


FIG. 3B

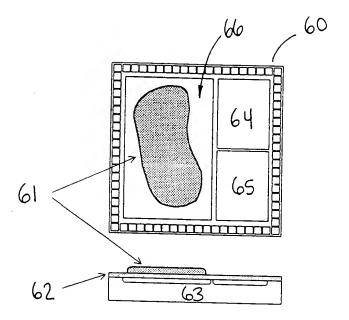


FIG. 4

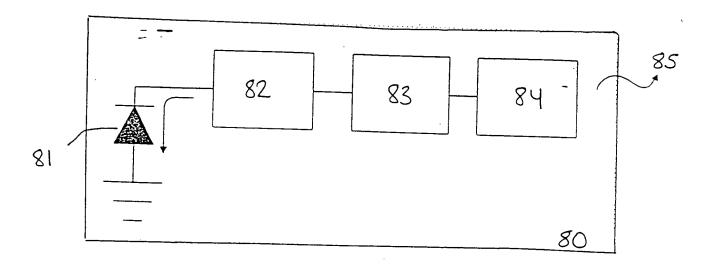


FIG. 5

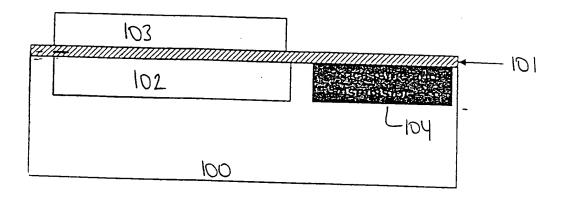


FIG. 6

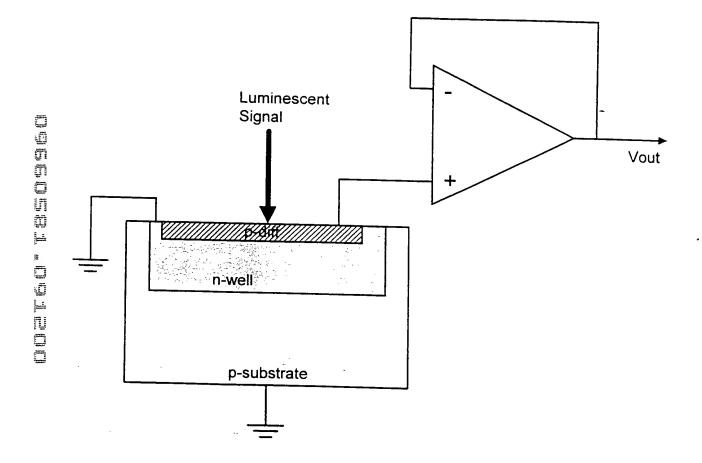


FIG. 7A

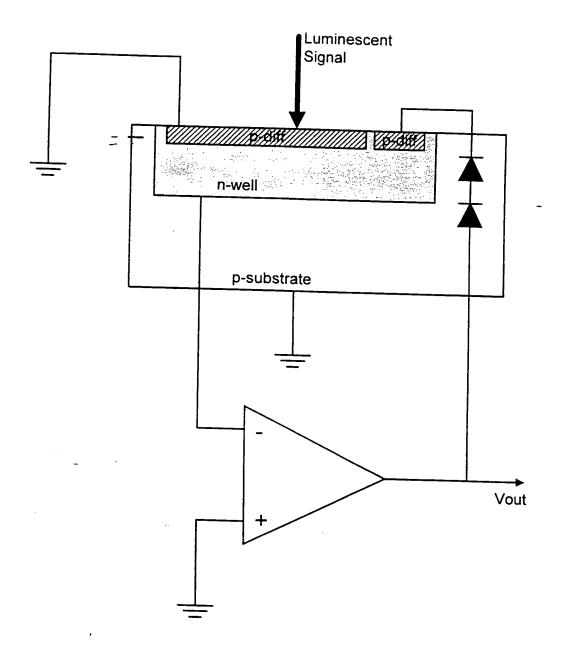


FIG. 7B

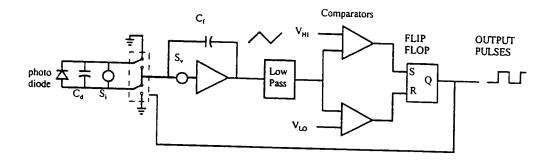


FIG. 7C

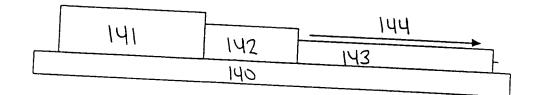


FIG. 8

FIG. 9A

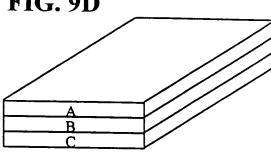
A	В	С
D	Е	F
G	Н	I
J	K	L

 \mathbf{A}

В

В D A G Н K E L





				Synthesized Prime	er	
S' Nhe I		I end of Tn	5			
GGGCGCTAGCGAA	ATGTTGACTGT	CTCTTCAT	CAGAMOTOTO	AATTCAGAAGAACTCG		
CCCGCTGTGCCTT	TACAACTTATG	AGTATGAG	AAGGAAAAG'	TTAAGTCTTCTTGAGC		
		** ****	*** *	5'		
				W IC C		
				Km ^r Gene Sequen	ce	
				•		
				Synthesized Primer		
T. D.	_			- j /		
	of Tn5	Xba I	Not I	<i>y</i> a.		
CGAATTETCACTED	TATACACAAGT	TCTAGATT	GCGGCCGCTT	GGTTAAAAAATGAGC		
GATAAGAAAACTAA	ATATTCCCTAA	AAGCGCTA * * * * * * * * * * * * * * * * * * *	AGCCGGATAA	CCAATTTTTTACTCG		
	· -	** A	• ••			
				Km ^r Gene Sequenc	e	
				-		
imers for pLJS						
and of philo						
			Synth	esized Primer		
5. BssH II	Spe I Xba I	Nh	e I Avr 🛚 🌶	Kpn I		
•	AACTAGTCTAG	ACTAAAGO	TACCCTACCC	TGGGAMGG		
GGTTCGCGCG	TTAATTGGAGT	GATTTCCC	TTGTTTTCGG:			
3'	* ** **	** * *1	**** *	5'		
	•					
pBScript KS Gene Seque	nce					
de la come peque	nce					
	••					
5'	***	***		3′		
TCCAATTAGTGÄGTGTGTATTACGCGCGCTCAC CACGAGGTTAAGCGGGATCGAGCATAATGCGCGCAGTG						
3' Sac I	Avr II Nh	ATCGAGCA A Y				
- -	AH II IYA	2 T	BssH II	5′		
			_	<u> </u>		
			Synthes	sized Primer		

FIG. 10

Denotes base pair mismatch

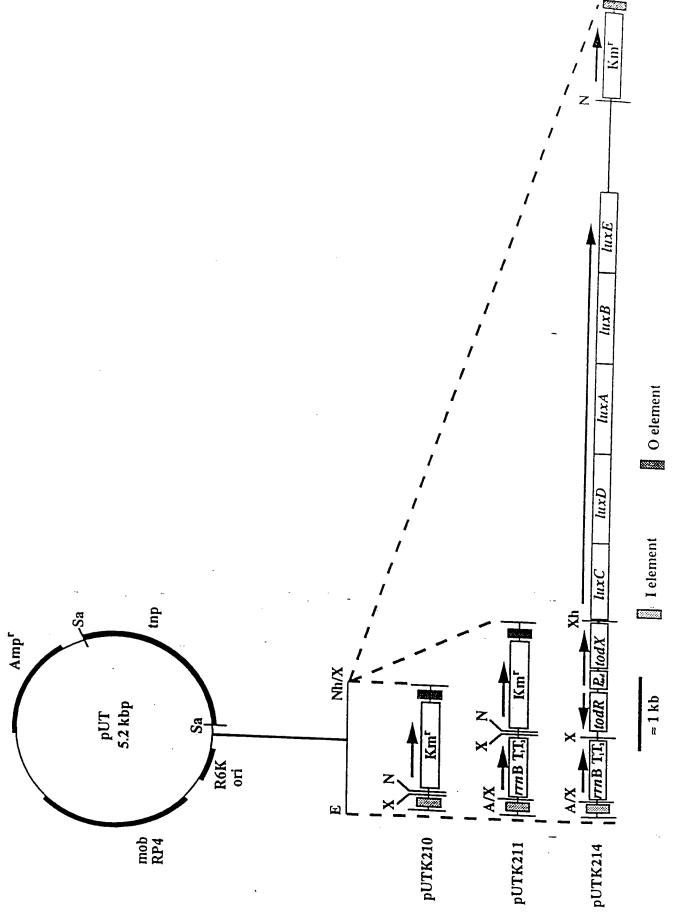


FIG. 11

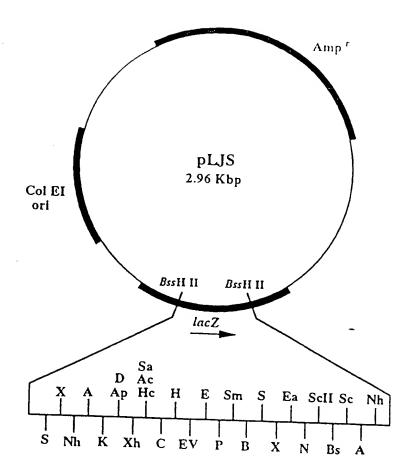


FIG. 12

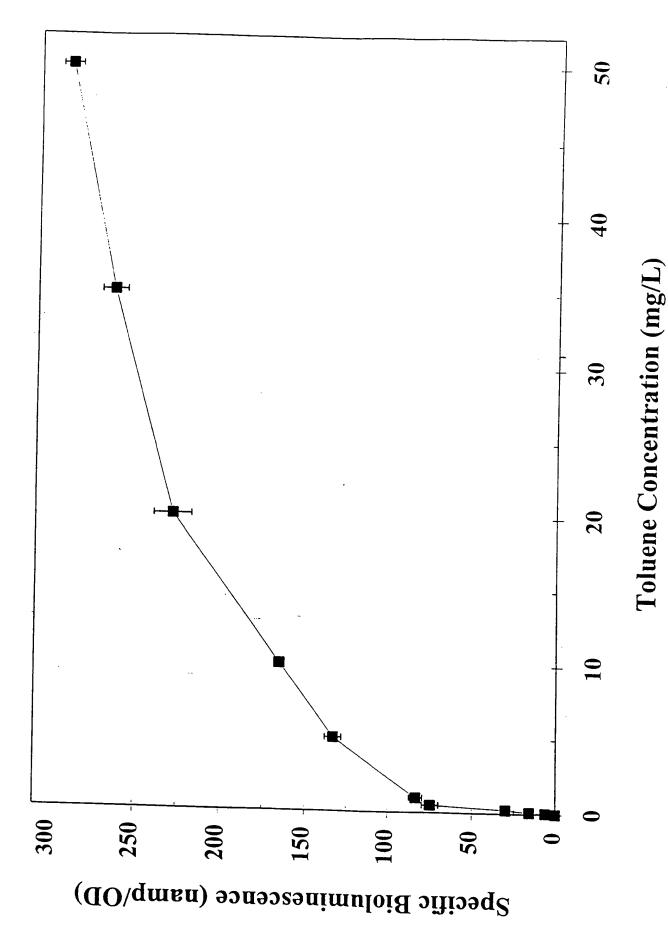
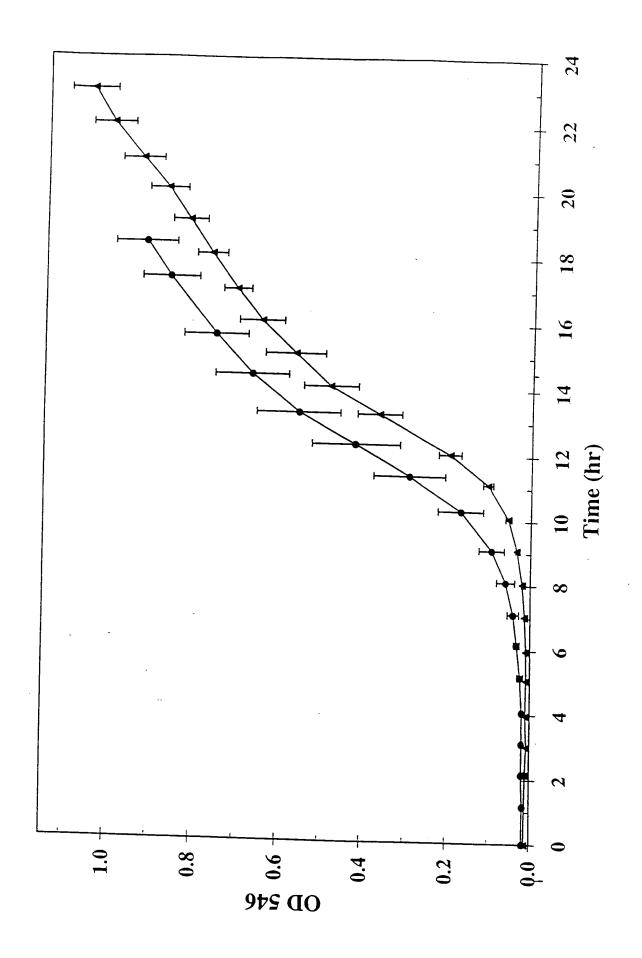
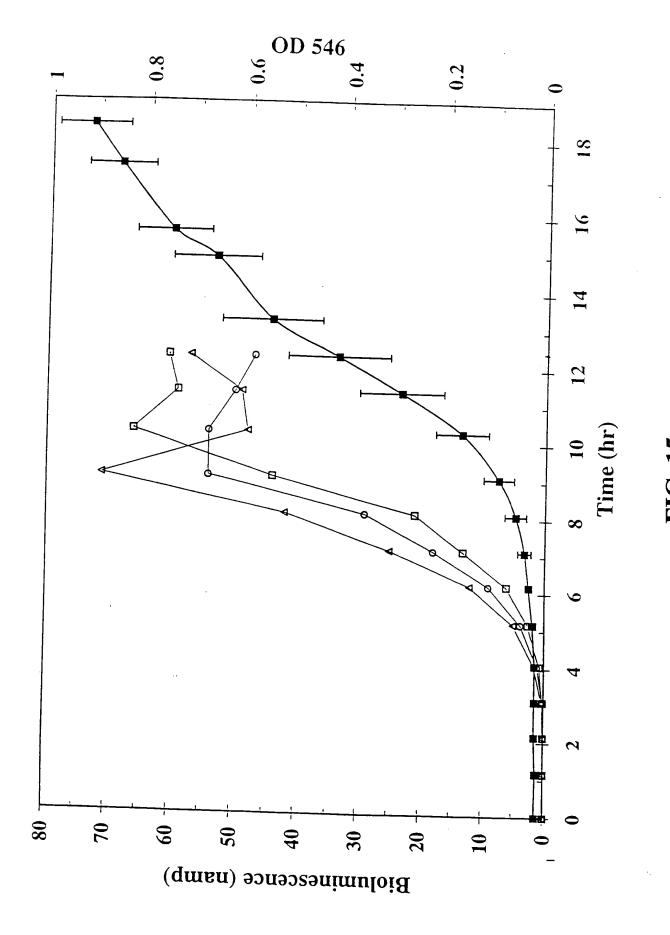


FIG. 13





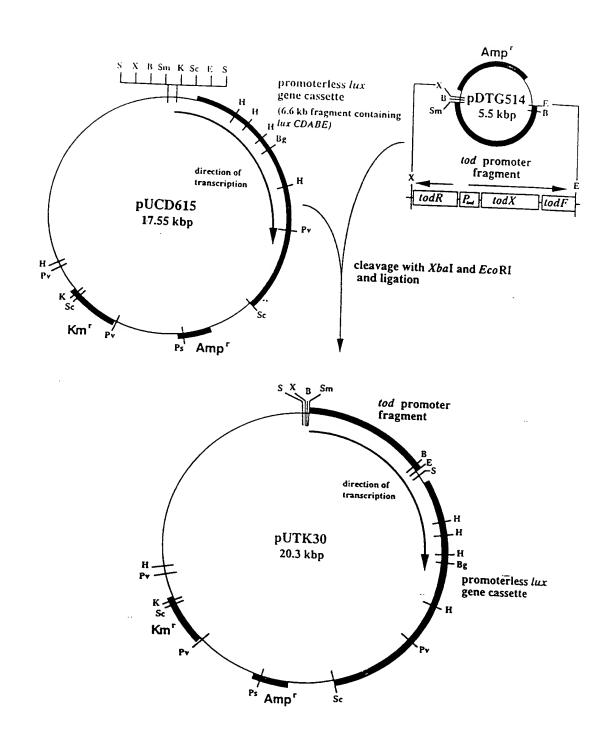


FIG. 16

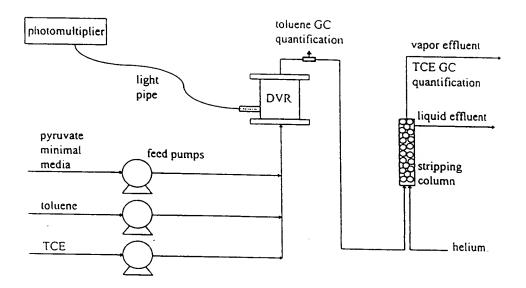


FIG. 17

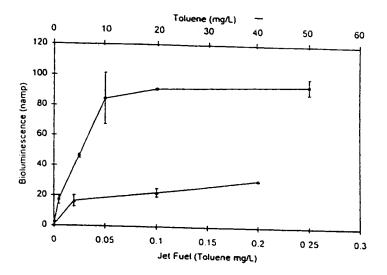


FIG. 18

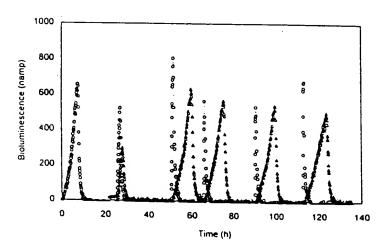


FIG. 19

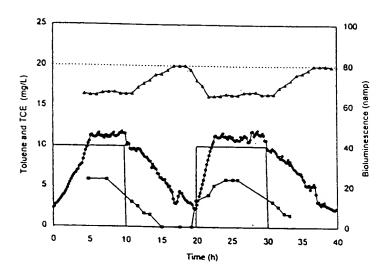


FIG. 20

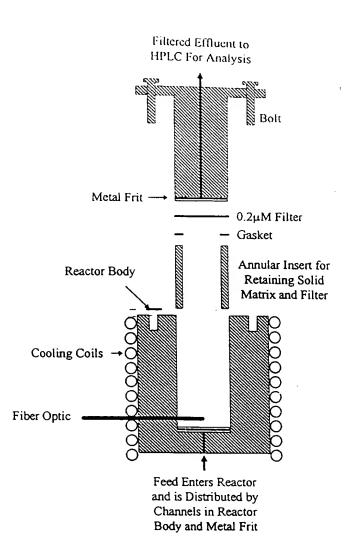


FIG. 21

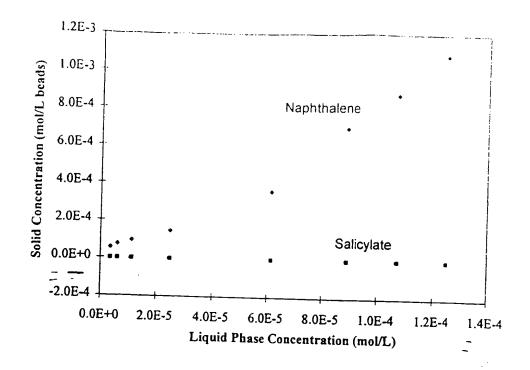


FIG. 22

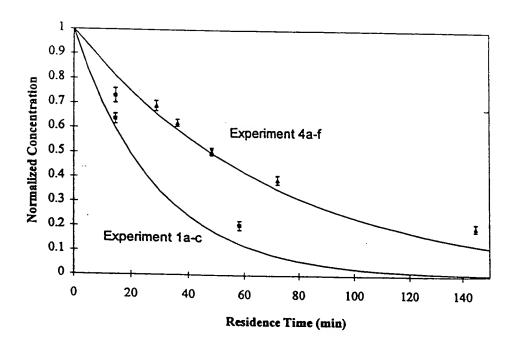


FIG. 23

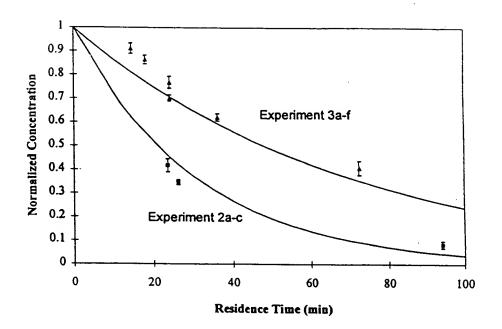


FIG. 24

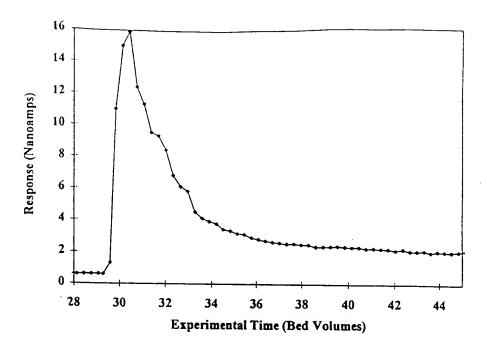


FIG. 25

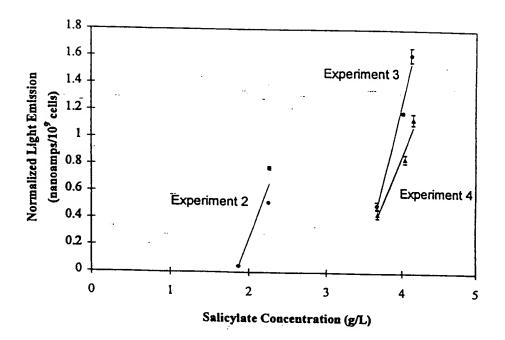


FIG. 26

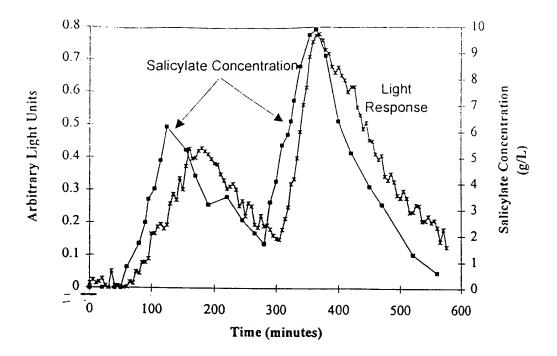


FIG. 27

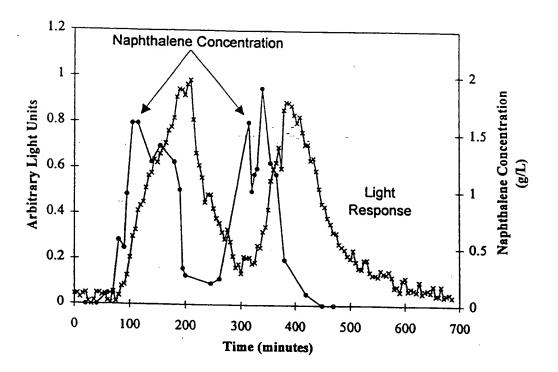


FIG. 28

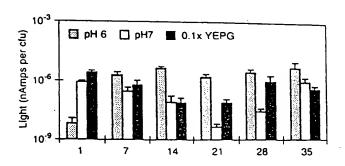


FIG. 29A

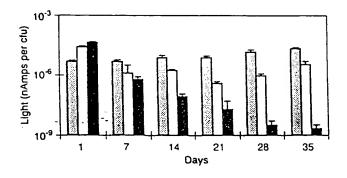


FIG. 29B

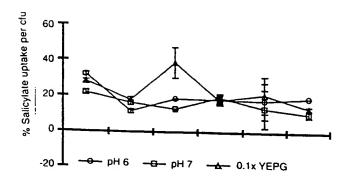


FIG. 30A

5

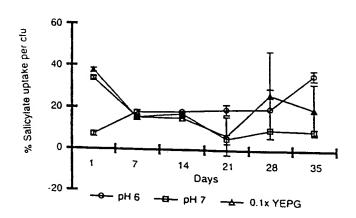
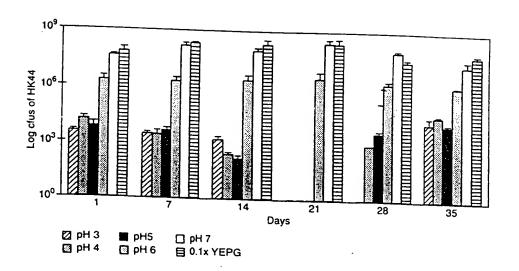


FIG. 30B

Fig. 3 Population of HK44 in alginate beads. The loganthm of the number of colony-forming units/alginate beads is shown



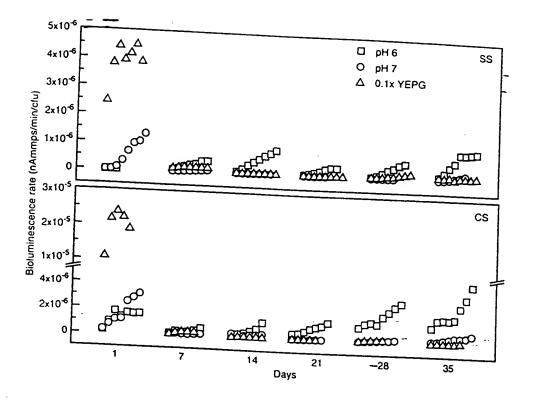


FIG. 32

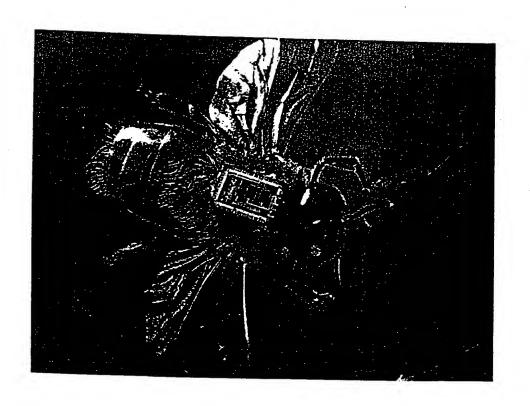


FIG. 33

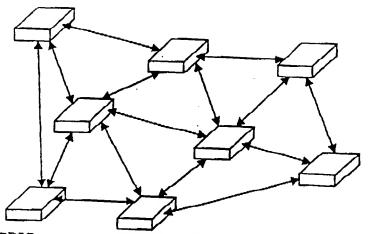
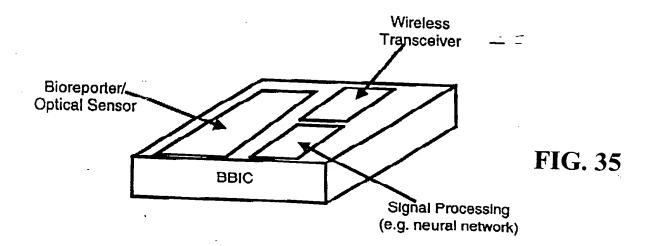
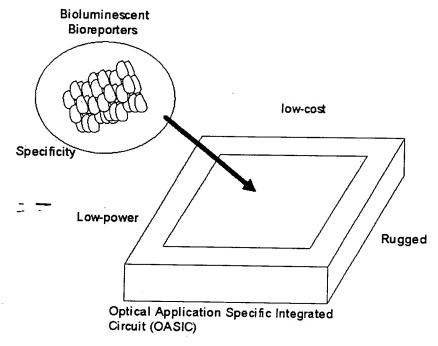


FIG. 34

BBICs connected together in a distributed neural network





High functional density
analog signal conditioning
digital signal processing
wireless transmission

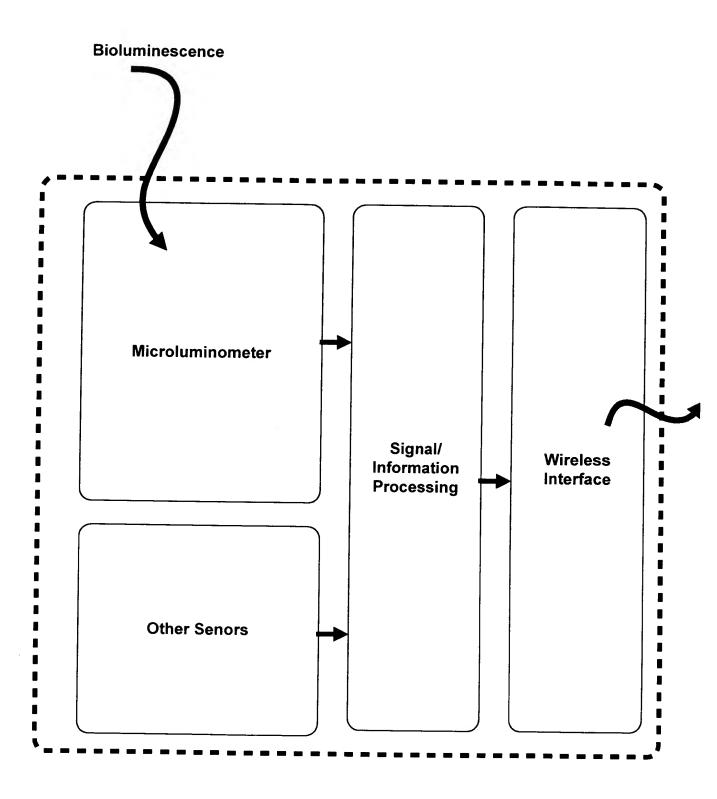
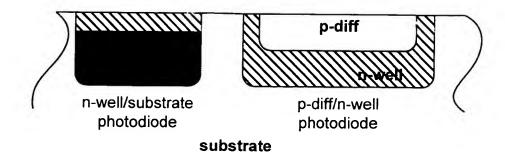


FIG. 37



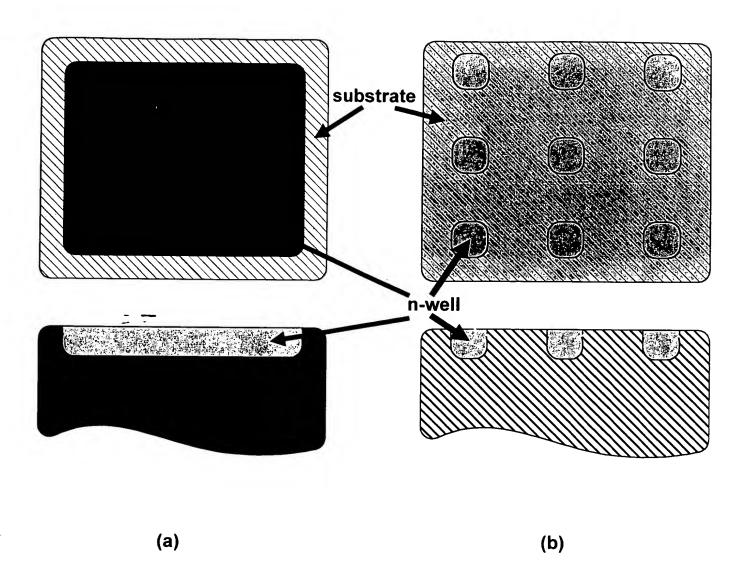


FIG. 39

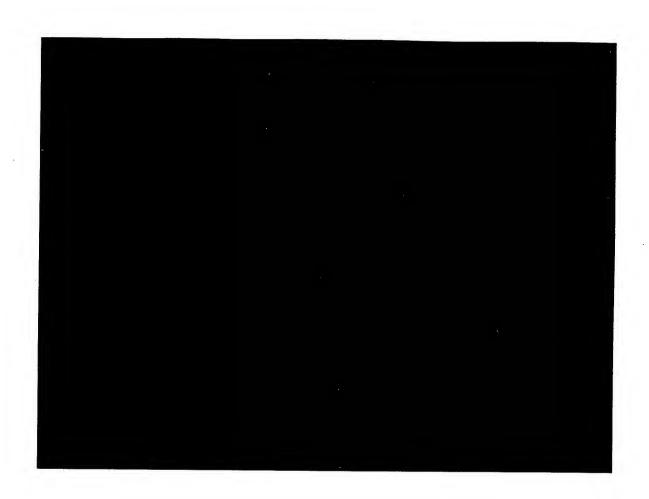


FIG. 40

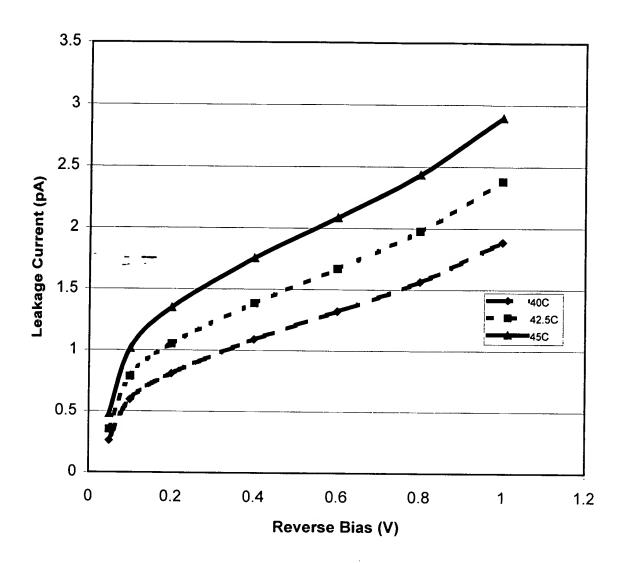


FIG. 41

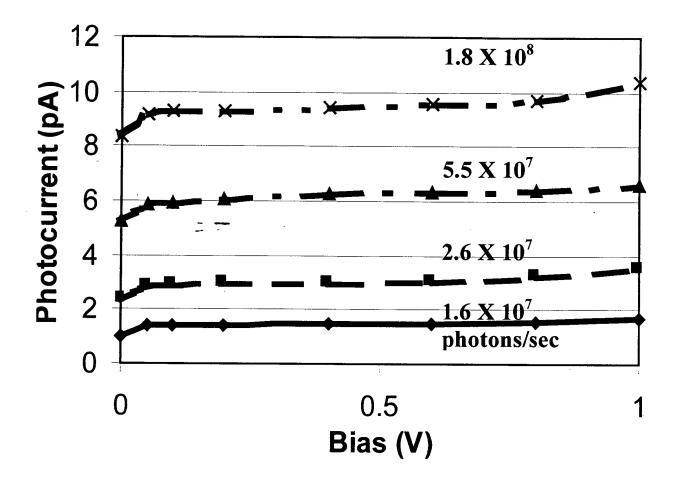


FIG. 42

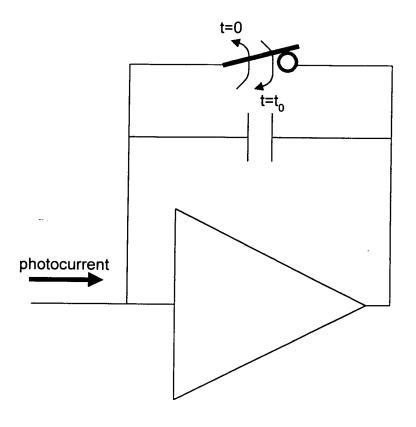


FIG. 43

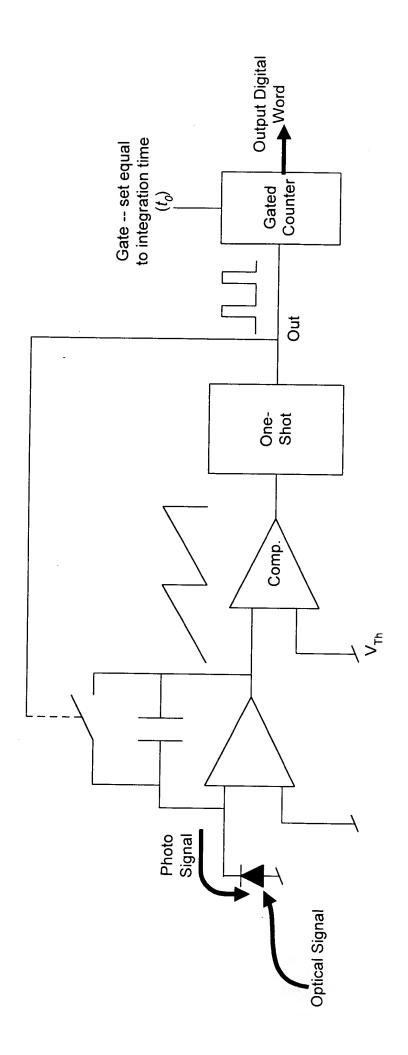


FIG. 44

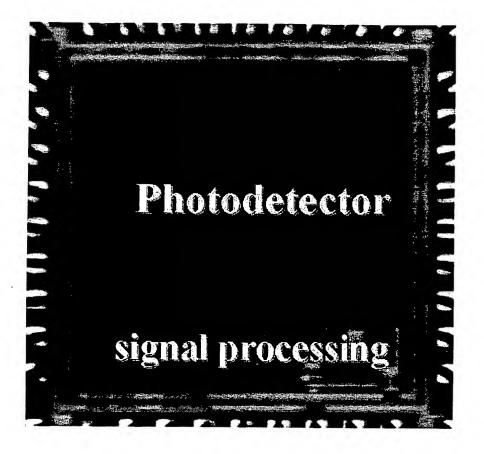


FIG. 45

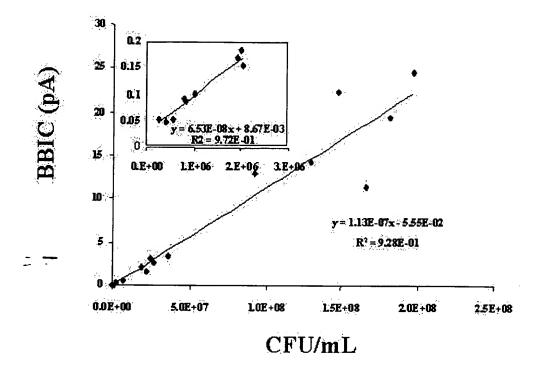


FIG. 46

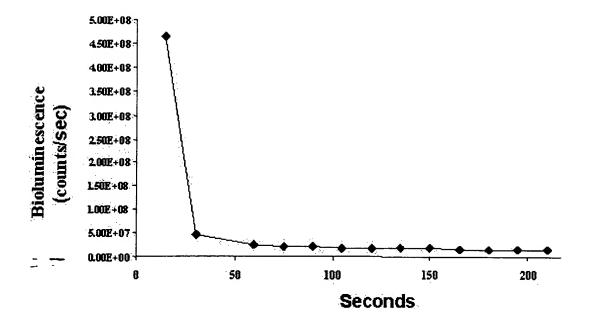
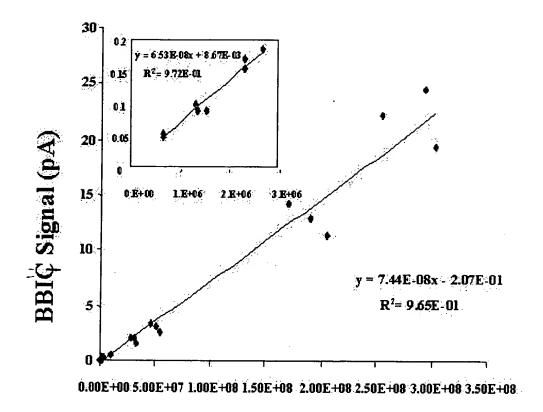


FIG. 47



Azur Luminometer (counts/s)

FIG. 48

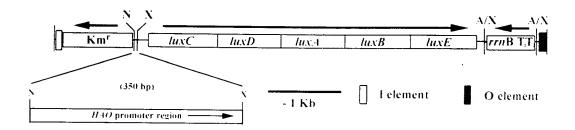
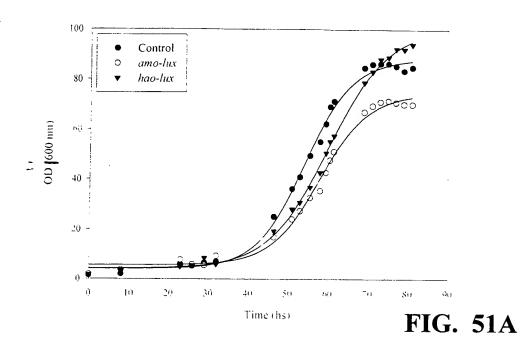


FIG. 49

	lux standards		samples		
	10 ng		control		hao3
1		0.1 ng		amo4	
	3 ng		amo1		hao4
	4	0.03 ng		aao1	
	l ng		amo2		pUTK-amo
	antest:				
		0.01 ng		hao2	
	0.3 ng		amo3		pUTK-hao

FIG. 50



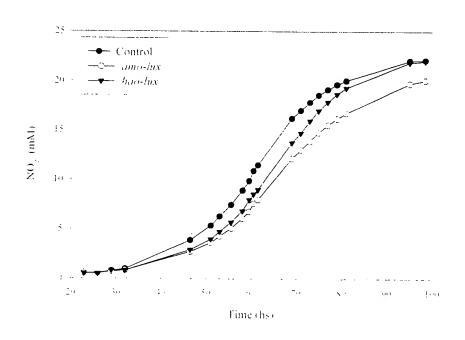


FIG. 51B

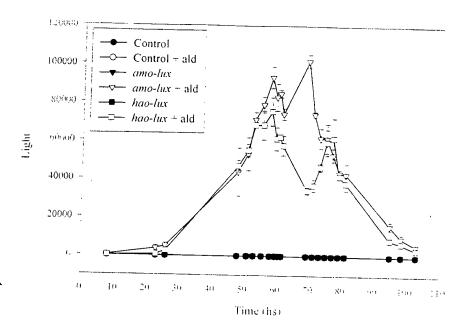


FIG. 52A

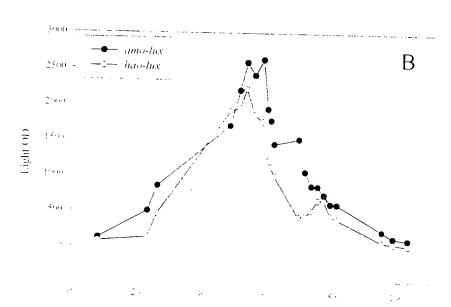


FIG. 52B

Lime that

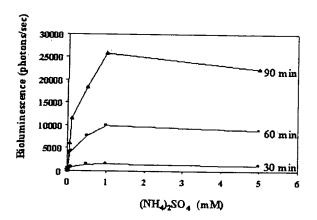


FIG. 53

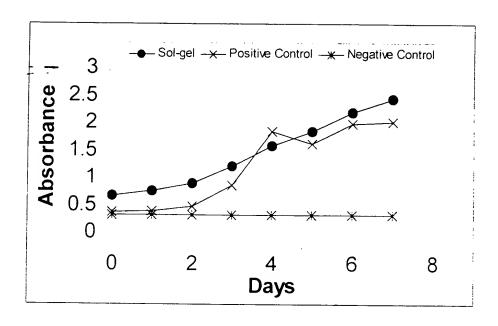


FIG. 54



FIG. 55

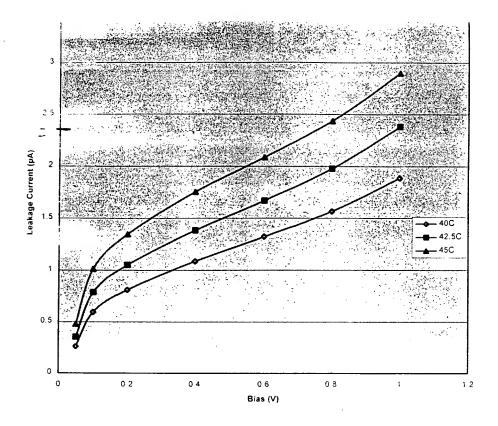


FIG. 56

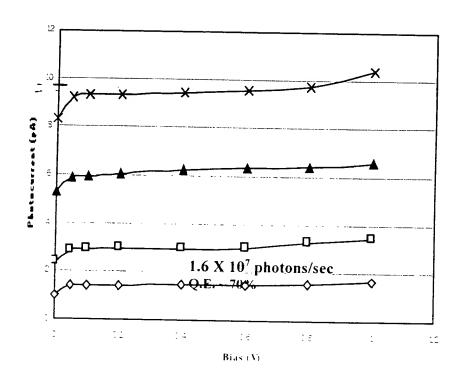


FIG. 57

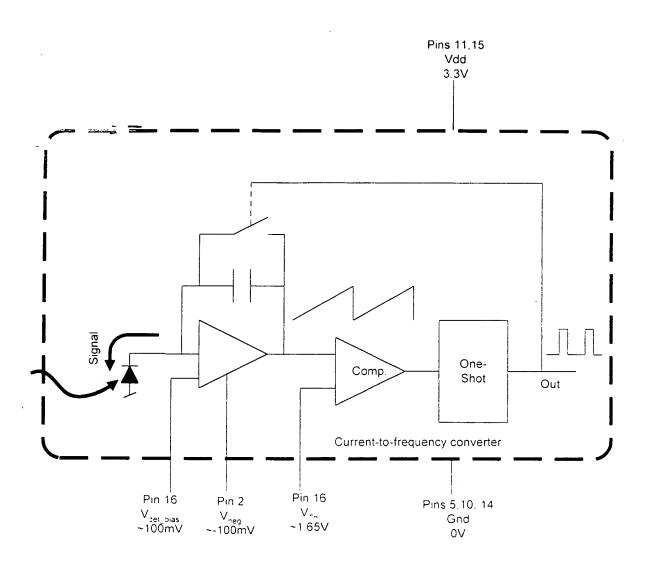


FIG. 58